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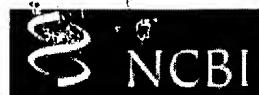
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Blast 2 Sequences results

PubMed

Entrez

BLAST

OMIM

Taxonomy

Structure

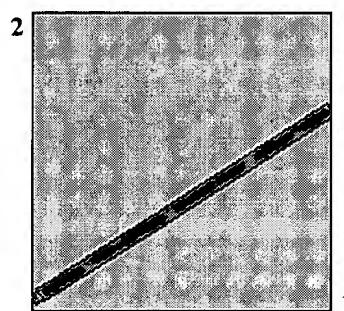
BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.5 [Nov-16-2002]

Match: 1 Mismatch: -2 gap open: 5 gap extension: 2

x dropoff: expect: wordsize: Filter Align

Sequence 1|seq_1 SEQ ID NO:5 Length 2222 (1 .. 2222)

Sequence gi_31543639 Homo sapiens solute carrier family 40 (iron-regulated transporter), member 1 (SLC40A1), mRNA Length 3460 (1..3460)



NOTE: The statistics (bitscore and expect value) is calculated based on the size of nr database.

NOTE: If protein translation is reversed, please repeat the search with reverse strand of the query sequence

Score = 4226 bits (2198), Expect = 0.0
Identities = 2211/2215 (99%), Gaps = 1/2215 (0%)
Strand = Plus / Plus

Query: 1 agctggctcagggcggtccgctaggctcgacgacctgctgagcctccaaaccgcttcca 60
Sbjct: 126 aactggactcagggcggtccgctaggctcgacgacctgctgagcctccaaaccgcttcca 185

Query: 61 taaggcttgc-tttccaacttcagctacagtgttagctaagttggaaagaaggaaaaaa 119
Sbjct: 186 taaggcttgccttccaacttcagctacagtgttagctaagttggaaagaaggaaaaaa 245

Query: 120 agaaaatccctggcccccctttctttgttcttgccaaagtgcgtgttagtctttt 179
Sbjct: 246 agaaaatccctggcccccctttctttgttcttgccaaagtgcgtgttagtctttt 305

Query: 180 gcccaaggctgtgttttagaggtgctatctccagttcctgcactcctgttaacaa 239
Sbjct: 306 gcccaaggctgtgttttagaggtgctatctccagttcctgcactcctgttaacaa 365

Query: 240 gcacacctcgagcgagcagcagcagcgatagcagccgcagaagagccagcgggtcgctc 299
Sbjct: 366 gcacacctcgagcgagcagcagcagcgatagcagccgcagaagagccagcgggtcgctgta 425

Query: 300 gtgtcatgaccaggcgaggatcacaaccgcacagagaggatgtggatcctggcg 359
Sbjct: 426 gtgtcatgaccaggcgaggatcacaaccgcacagagaggatgtggatcctggcg 485

Sbjct: 11 tt aa aga aggg aa act ga att gaa ac a g c t g a a c a c a a g a t a c t g a g c c a a a c 1205
solute carrier family 40 (iron-> 239 L K E E E T E L K Q L N L H K D T E P K

Query: 1080 cc ct gg ag gg a a c t c a t c t a a t g g g t g a a a g a c t c t a a c a t c c a t g a g c t t g a a c a t g 1139
Sbjct: 1206 cc ct gg ag gg a a c t c a t c t a a t g g g t g a a a g a c t c t a a c a t c c a t g a g c t t g a a c a t g 1265
solute carrier family 40 (iron-> 259 P L E G T H L M G V K D S N I H E L E H

Query: 1140 a g c a a g a g c c t a c t t g t g c c t c c a g a t g g c t g a g c c t t c c g t a c c t c c g a g a t g g a t 1199
Sbjct: 1266 a g c a a g a g c c t a c t t g t g c c t c c a g a t g g c t g a g c c t t c c g t a c c t c c g a g a t g g a t 1325
solute carrier family 40 (iron-> 279 E Q E P T C A S Q M A E P F R T F R D G

Query: 1200 g g g t c t c t a c t a c a a c c a c c a g c c t g t g t t c t g g c t g g c a t g g g t c t t g c t t c t t t a t a 1259
Sbjct: 1326 g g g t c t c t a c t a c a a c c a c c a g c c t g t g t t c t g g c t g g c a t g g g t c t t g c t t c t t t a t a 1385
solute carrier family 40 (iron-> 299 W V S Y Y N Q P V F L A G M G L A F L Y

Query: 1260 t g a c t g t c c t g g g c t t g a c t g c a t c a c c a c a g g g t a c g c c t a c a c t c a g g g a c t g a g t g 1319
Sbjct: 1386 t g a c t g t c c t g g g c t t g a c t g c a t c a c c a c a g g g t a c g c c t a c a c t c a g g g a c t g a g t g 1445
solute carrier family 40 (iron-> 319 M T V L G F D C I T T G Y A Y T Q G L S

Query: 1320 g t t c c a t c c t c a g t a t t t g a t g g g a g c a t c a g c t a t a c t g g a a t a t g g g a a c t g t a g 1379
Sbjct: 1446 g t t c c a t c c t c a g t a t t t g a t g g g a g c a t c a g c t a t a c t g g a a t a t g g g a a c t g t a g 1505
solute carrier family 40 (iron-> 339 G S I L S I L M G A S A I T G I M G T V

Query: 1380 c t t t a c t t g g c t a c g t c g a a a t g t g g t t g g t c g a c a g g t c t g a t c t c a g g a t t g g 1439
Sbjct: 1506 c t t t a c t t g g c t a c g t c g a a a t g t g g t t g g t c g a c a g g t c t g a t c t c a g g a t t g g 1565
solute carrier family 40 (iron-> 359 A F T W L R R K C G L V R T G L I S G L

Query: 1440 c a c a g c t t c c t g t t g a t c t t g t g t g a t c t c t g t a t t c a t g c c t g g a a g c c c c t g g 1499
Sbjct: 1566 c a c a g c t t c c t g t t g a t c t t g t g t g a t c t c t g t a t t c a t g c c t g g a a g c c c c t g g 1625
solute carrier family 40 (iron-> 379 A Q L S C L I L C V I S V F M P G S P L

Query: 1500 a c t t g t c c g t t t c t c t t t g a a g a t a t c c g a t c a a g g t t c a t t c a a g g a g a g t c a a t t a 1559
Sbjct: 1626 a c t t g t c c g t t t c t c t t t g a a g a t a t c c g a t c a a g g t t c a t t c a a g g a g a g t c a a t t a 1685
solute carrier family 40 (iron-> 399 D L S V S P F E D I R S R F I Q G E S I

Query: 1560 c a c c t a c c a a g a t a c c t g a a a t t a c a a c t g a a a t a t a c a t g t c t a a t g g g t c t a a t t c t g 1619
Sbjct: 1686 c a c c t a c c a a g a t a c c t g a a a t t a c a a c t g a a a t a t a c a t g t c t a a t g g g t c t a a t t c t g 1745
solute carrier family 40 (iron-> 419 T P T K I P E I T T E I Y M S N G S N S

Query: 1620 c t a a t a t t g t c c c g g a g a c a a g t c c t g a a t c t g t g c c c t a a t c t c t g t c a g t c t g c t g t 1679
Sbjct: 1746 c t a a t a t t g t c c c g g a g a c a a g t c c t g a a t c t g t g c c c t a a t c t c t g t c a g t c t g c t g t 1805
solute carrier family 40 (iron-> 439 A N I V P E T S P E S V P I I S V S L L

Query: 1680 t t g c a g g c g t c a t t g c t g c t g a a t c g g t c t t g g t c t t g a t t a c t g t g a c a c a g t 1739
Sbjct: 1806 t t g c a g g c g t c a t t g c t g c t g a a t c g g t c t t g g t c t t g a t t a c t g t g a c a c a g t 1865
solute carrier family 40 (iron-> 459 F A G V I A A R I G L W S F D L T V T Q

Query: 1740 t g c t g c a a g a a a t g t a a t g a a t c t g a a a g a g g c a t t a a t g g t g t a c a g a a c t c c a 1799

Blast Result

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T: 0
A: 0

X1: 6 (11.5 bits)
X2: 26 (50.0 bits)
S1: 12 (23.8 bits)
S2: 21 (41.1 bits)